- a) said homology is at least about 90% identity and said portion is at least about 9 amino acids;
- b) said homology is at least about 80% identity and] said [portion] sequence of 100% identity is over at least [about] 17 contiguous amino acids[; or
- c) said homology is at least about 70% identity and said portion is at least about 25 amino acids].
- 3. The composition of matter of Claim 1, wherein\said:
- a) 499E9 comprises a mature sequence of Table 1 (see SEO ID NO: 1); or
  - b) protein or peptide [:

5

15

20

25

- i) ] is from a (warm blooded animal selected from a)
   mammal [, including a rodent;
- ii) comprises at least one polypeptide segment of SEQID NO: 2;
- iii) exhibits a plurality of portions exhibiting said identity;
- iv) is a natural allelic variant of 499E9;
- v) has a length at least about 30 amino acids;
- vi) exhibits at least two non-overlapping epitopes which are specific for a mammalian 499E9;
- vii) exhibits a sequence identity at least about 90% over a length of at least about 20 amino acids to a rodent 499E9;
- viii) exhibits at least two non-overlapping epitopes which are specific for a rodent 499E9;
- ix) exhibits a sequence identity at least about 90%
   over a length of at least about 20 amino acids to a
   rodent 499E9;
- x) is glycosylated;
- xi) is a synthetic polypeptide;
- xii) is attached to a solid substrate;
- xiii) is conjugated to another chemical moiety;
- 35 xiv) is a 5-fold or less substitution from natural sequence; or

- xv) is a deletion or insertion variant from a natural sequence].
- 4. A composition of matter of Claim 1 which is sterile
  5 [comprising:
  - a) a sterile 499E9 protein or peptide of Claim 1; or
  - b) said 499E9 protein or peptide of Claim 1 and a carrier, wherein said carrier is:
    - i) an aqueous compound, including water, saline, and/or buffer; and/or
    - ii) formulated for oral, rectal, nasal, topical, or parenteral administration).
  - 5. The fusion protein of Claim 1, comprising:
- a) mature protein comprising sequence of Table 1 <u>(see SEO ID NO: 2)</u>;
  - a detection or purification tag, including a FLAG, His6, or Ig sequence; or
  - c) sequence of another [TNF] <u>tumor necrosis factor</u> ligand protein.
  - 6. A kit comprising a [protein or polypeptide of Claim 1, and:
    - a) a] compartment comprising said [protein or] polypeptide of Claim 1 [; and/or
    - b) ] and instructions for use or disposal of reagents in said kit.
- 11. An isolated or recombinant nucleic acid encoding a [protein or peptide] polypeptide or fusion protein of Claim 1, wherein [:
  - a) ] said 499E9 protein is from a mammal [, including a rodent; or
  - b) said nucleic acid:
    - i) encodes an antigenic peptide sequence of Table 1;
    - ii) encodes a plurality of antigenic peptide sequences of Table 1;

10

20

25

- iii) exhibits at least about 80% identity to a natural cDNA encoding said segment; is an expression vector; v) further comprises an origin of replication; is from a natural source; vii) comprises a detectable label; viii) comprises synthetic nucleotide sequence; ix) is less than 6 kb, preferably less than 3 kb;
  - is from a mammal, including a rodent;
  - comprises a natural full length coding sequence; xii) is a hybridization probe for a gene encoding said TNF-ligand family protein; or
  - xiii) is a PCR primer, PCR product, or mutagenesis primer].

14. A kit comprising [said nucleic acid of Claim 11, and:

- a) ] a compartment comprising said nucleic acid of Claim 11 [;
- b) a compartment further comprising a 499E9 protein or polypeptide; and/or
- c) ] and instructions for use or disposal of reagents in said kit.
- 15. A nucleic acid which [:
- 25 a) ] selectively hybridizes under wash conditions of [30] at least 45° C and less than [2M] 500 mM salt to SEQ ID NO: 1[; or
  - exhibits at least about 85% identity over a stretch of at b) least about 30 nucleotides to a rodent 499E9].
  - 16. The nucleic acid of Claim 15, wherein:
    - said wash conditions are at [45] <u>least 55° C [and/or 500]</u> and less than 150 mM salt; or

DEC 17 '98 17:00 FR DNAX RESEARCH INST. 415 496 1200 TO 917033057401

said [identity is at least 90% and/or said stretch is] b) 35 nucleic acid comprises at least [55] 30 contiguous nucleotides of the coding portion of SEO ID NO: 1.

5

10

15

20

## Please add new Claims 21-46 as follows:

- --21. The composition of matter of Claim 1, which comprises the natural sequence 499E9 of SEQ ID NO: 2.
- 22. The recombinant 499E9 polypeptide of Claim 2, wherein said 100% identity is over at least 25 contiguous amino acids.
- 23. The substantially pure 499E9 polypeptide of Claim 2, wherein said 100% identity is over at least 30 contiguous amino 10 acids.
  - 24. The substantially pure 499E9 polypeptide of Claim 1, which has a length of at least 30 amino acids.
  - 25. The substantially pure or recombinant 499E9 polypeptide of Claim 1, which is:
    - a) glycosylated;

5

15

25

35

- a synthetic polypeptide;
- 20 attached to a solid substrate; or
  - conjugated to another chemical entity.
  - 26. A composition comprising said 499E9 polypeptide of Claim 1 and an aqueous carrier.
  - 27. The composition of Claim 26, formulated for oral, rectal, nasal, topical, or parenteral administration.
- 28. The nucleic acid of Claim 11, which comprises at least 30 22 contiguous nucleotides of the coding portion of SEQ ID NO: 1.
  - 29. An isolated or recombinant nucleic acid which encodes a polypeptide or fusion protein of Claim 1, wherein said polypeptide is an antigenic peptide of Table 1 (see SEQ ID NO: 2).
  - 30. The nucleic acid of Claim 29, which comprises at least 29 contiguous nucleotides of the coding portion of SEQ ID NO: 1.

GORMAN and MATTSON; USSN 08/989,362

- 31. An isolated or recombinant nucleic acid encoding a polypeptide of Claim 1, which exhibits 100% identity over the mature protein coding portion to a natural DNA encoding said 499E9.
- 32. A vector which encodes a 499E9 polypeptide of Claim 1 and comprises:
  - at least 35 contiguous nucleotides of the coding portion of SEQ ID NO: 1;
  - b) transcriptional regulatory sequences operably linked to said 499E9 coding sequence; or
  - an origin of replication. C)
- 15 33. The vector of Claim 32, comprising at least 41 contiguous nucleotides from the coding portion of SEQ ID NO: 1.
- 34. An isolated or recombinant nucleic acid encoding a polypeptide or fusion protein of Claim 1, wherein said nucleic 20 acid:
  - a) is from a natural source;
  - comprises a detectable label;
  - comprises synthetic nucleotide sequence; or
  - d) comprises natural full length coding sequence.
  - 35. An isolated or recombinant nucleic encoding a polypeptide of Claim 1, which is a hybridization probe for a gene encoding a tumor necrosis factor ligand family protein.
- 30 36. A cell comprising said nucleic acid of Claim 29.
  - 37. A cell comprising said nucleic acid of Claim 31.
  - 38. A cell comprising said nucleic acid of Claim 32.
    - 39. A cell comprising said nucleic acid of Claim 34.

GORMAN and MATTSON; USSN 08/989,362

5

10

25

- 40. A kit comprising a compartment comprising a nucleic acid of Claim 34 and instructions for use or disposal of reagents in said kit.
- 5 A kit comprising a compartment comprising said nucleic acid of Claim 35 and instructions for use or disposal of reagents in said kit.
- 42. A method of making a protein, comprising culturing a 10 cell of Claim 12 in an environment resulting in expressing said protein and recovering said protein.
  - 43. A method of making a protein, comprising culturing a cell of Claim 29 in an environment resulting in expressing said protein and recovering said protein.
  - 44. A method of making a protein, comprising culturing a cell of Claim 32 in an environment resulting in expressing said protein and recovering said protein.
  - 45. A method of making a duplex nucleic acid comprising contacting a nucleic acid of Claim 29 with a complementary nucleic acid under selective hybridization conditions of at least 45° C and less than 500 mM salt, thereby forming said duplex.
  - A method of making a polynucleotide of Claim 11, comprising amplifying said polypeptide using PCR amplification methods. --

GORMAN and MATTSON; USSN 08/989,362

15

20